Figure 1A

ATGTCGAAAA TTGAACTAAA ACAACTATCT TTTGCCTATG ATAATCAAGA AGTATTGCTT 60
TTTGATCAGG CAAATATCAC GATGGATACC AATTGGAAAT TAGGATTGAT TGGCCGCAAT 120
GGCCGTGGGA AAACAACCTT ATTAAGATTG TTACAAAAAC AGTTGGATTA CCAAGGAGAG 180
ATTCTTCATC AAGTCGATTT CGTCTATTTT CCACAAACAG TTGCAGAAGA ACAACAGCTC 240
ACTTATTATG TCTTACAAGA GGTGACTTCT TTTGAACAGT GGGAATTAGA ACGAGAATTA 300
ACGCTTTTAA ACGTTGATCC TGAAGTTTTA TGGCGGCCCT TTTCTTCTTT ATCAGGCGGC 360
GAAAAGACGA AAGTTTTATT AGGTCTTCTT TTTATTGAAG AAAATGCCTT TCCTTTAATT 420
GACGAGCCAA CAAATCATTT AGATCTAGCT GGCAGACAAC AAGTGGCTGA ATATTTGAAG 480
AAAAAGAAAC ACGGGTTTAT TTTAGTCAGC CACGATCGGG CATTTGTTGA TGAAGTGGTT 540
GATCATATTT TGGCGATTGA AAAAAGTCAA TTGACGCTGT ATCAAGGGAA TTTTTCTATT 600
TATGAAGAGC AAAAAAAATT AAGAGATGCT TTTGAACTAG CAGAAAATGA AAAAATCAAA 660
AAAGAAGTCA ATCGCTTGAA AGAAACCGCT CGTAAAAAAG CGGAATGGTC GATGAACCGT 720
GAAGGTGATA AGTACGGCAA CGCTAAGGAA AAAGGGAGCG GGCGATTTTT TGATACAGGA 780
GCCATTGGTG CCCGGGCAGC GCGCGTAATG AAGCGCTCGA AACACATTCA ACAACGCGCC 840
GAAACACAAT TAGCAGAAAA AGAAAAACTA TTAAAAGATC TTGAGTATAT TGATCCTTTG 900
TCAATGGATT ATCAGCCAAC GCATCACAAA ACATTATTGA CGGTGGAAGA GCTTCGTCTA 960

Figure 1B

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GGCTACGAGA AAAATTGGCT ATTTGCGCCA CTTTCTTTTT CAATAAACGC GGGAGAAATT  1020
GTTGGAATAA CAGGGAAAAA TGGCTCAGGA AAATCGAGCT TAATTCAGTA TTTATTGGAT  1080
AATTTTTCTG GGGATTCAGA AGGCGAAGCC ACTTTGGCTC ACCAATTAAC CATTTCTTAT  1140
GTGCGCCAAG ATTATGAAGA CAATCAAGGA ACTTTATCCG AATTTGCAGA GAAAAATCAG  1200
TTAGATTACA CTCAATTTTT AAATAACTTA CGAAAACTTG GGATGGAGCG CGCCGTTTTC  1260
ACTAATCGAA TTGAACAAAT GAGTATGGGG CAACGGAAAA AAGTCGAAGT AGCCAAATCA  1320
TTGTCTCAAT CAGCTGAACT TTATATTTGG GATGAACCCC TTAATTACTT GGATGTATTT  1380
AATCATCAAC AATTAGAAGC GCTAATCTTA TCTGTGAAGC CTGCAATGCT AGTGATTGAG  1440
CATGATGCAC ATTTCATGAA GAAAATAACA GATAAAAAAA TTGTCTTGAA ATCATAA     1497
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Figure 2A

MetSerLysIleGluLeuLysGlnLeuSerPheAlaTyrAspAsnGlnGluValLeuLeu 20

PheAspGlnAlaAsnIleThrMetAspThrAsnTrpLysLeuGlyLeuIleGlyArgAsn 40

GlyArgGlyLysThrThrLeuLeuArgLeuLeuGlnLysGlnLeuAspTyrGlnGlyGlu 60

IleLeuHisGlnValAspPheValTyrPheProGlnThrValAlaGluGluGlnGlnLeu 80

ThrTyrTyrValLeuGlnGluValThrSerPheGluGlnTrpGluLeuGluArgGluLeu 100

ThrLeuLeuAsnValAspProGluValLeuTrpArgProPheSerSerLeuSerGlyGly 120

GluLysThrLysValLeuLeuGlyLeuLeuPheIleGluGluAsnAlaPheProLeuIle 140

AspGluProThrAsnHisLeuAspLeuAlaGlyArgGlnGlnValAlaGluTyrLeuLys 160

LysLysLysHisGlyPheIleLeuValSerHisAspArgAlaPheValAspGluValVal 180

AspHisIleLeuAlaIleGluLysSerGlnLeuThrLeuTyrGlnGlyAsnPheSerIle 200

TyrGluGluGlnLysLysLeuArgAspAlaPheGluLeuAlaGluAsnGluLysIleLys 220

LysGluValAsnArgLeuLysGluThrAlaArgLysLysAlaGluTrpSerMetAsnArg 240

GluGlyAspLysTyrGlyAsnAlaLysGluLysGlySerGlyAlaIlePheAspThrGly 260

AlaIleGlyAlaArgAlaAlaArgValMetLysArgSerLysHisIleGlnGlnArgAla 280

GluThrGlnLeuAlaGluLysGluLysLeuLeuLysAspLeuGluTyrIleAspProLeu 300

SerMetAspTyrGlnProThrHisHisLysThrLeuLeuThrValGluGluLeuArgLeu 320

Figure 2B

GlyTyrGluLysAsnTrpLeuPheAlaProLeuSerPheSerIleAsnAlaGlyGluIle 340

ValGlyIleThr<u>GlyLysAsnGlySerGlyLysSerSerLeu</u>IleGlnTyrLeuLeuAsp 360

AsnPheSerGlyAspSerGluGlyGluAlaThrLeuAlaHisGlnLeuThrIleSerTyr 380

ValArgGlnAspTyrGluAspAsnGlnGlyThrLeuSerGluPheAlaGluLysAsnGln 400

LeuAspTyrThrGlnPheLeuAsnAsnLeuArgLysLeuGlyMetGluArgAlaValPhe 420

ThrAsnArgIleGluGlnMetSerMetGlyGlnArgLysLysValGluValAlaLysSer 440

LeuSerGlnSerAlaGluLeuTyrIleTrp<u>AspGluProLeuAsnTyrLeuAsp</u>ValPhe 460

AsnHisGlnGlnLeuGluAlaLeuIleLeuSerValLysProAlaMetLeuValIleGlu 480

HisAspAlaHisPheMetLysLysIleThrAspLysLysIleValLeuLysSer 498

Figure 3A

ATGAAAGAGA	TCGTAACATT	AACAAACGTT	AGCTATGAAG	TAAAGGATCA	AACTGTTTTT	60
AAACATGTAA	ACGCCAGTGT	TCAGCAAGGA	GATATCATTG	GGATTATCGG	CAAAAACGGC	120
GCTGGGAAAT	CTACGTTGCT	GCACCTCATT	CACAATGACT	TAGCCCCTGC	ACAGGGTCAA	180
ATCCTTCGGA	AGGATATAAA	ACTGGCTTTG	GTTGAACAGG	AAACCGCGGC	GTATTCCTTT	240
GCGGATCAGA	CACCTGCCGA	AAAGAAGTTA	CTGGAGAAAT	GGCATGTGCC	TCTTCGTGAT	300
TTTCATCAGT	TAAGCGGCGG	TGAAAAACTG	AAAGCGCGGC	TGGCGAAAGG	ACTATCAGAG	360
GATGCAGATC	TGCTGCTGTT	AGATGAACCG	ACAAACCACC	TTGATGAAAA	AAGCTTGCAA	420
TTTCTCATCC	AACAGCTGAA	ACATTATAAC	GGCACTGTGA	TTCTCGTTTC	TCACGATCGA	480
TATTTTTTAG	ACGAAGCCGC	AACAAAAATA	TGGTCGCTTG	AGGATCAGAC	GCTGATTGAA	540
TTCAAAGGGA	ATTACTCCGG	GTATATGAAG	TTCCGGGAGA	AGAAAAGACT	CACCCAGCAG	600
CGTGAATATG	AAAAGCAGCA	AAAAATGGTT	GAACGGATTG	AAGCACAAAT	GAATGGGCTC	660
GCTTCTTGGT	CGGAAAAAGC	CCATGCTCAA	TCGACGAAAA	AGGAAGGGTT	TAAAGAATAT	720
CACCGGGTAA	AAGCGAAGCG	TACGGATGCC	CAGATAAAAT	CCAAGCAGAA	GCGGCTTGAA	780
AAAGAGCTTG	AAAAAGCAAA	GGCGGAACCC	GTTACCCCAG	AATATACAGT	CCGCTTTTCA	840
ATCGATACAA	CCCACAAAAC	AGGAAAACGT	TTTTTAGAAG	TTCAGAATGT	AACAAAAGCG	900
TTTGGAGAAA	GGACTCTCTT	TAAAAACGCA	AACTTTACAA	TTCAGCACGG	CGAAAAGGTT	960

Figure 3B

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GCGATCATAG GCCCCAATGG CAGCGGAAAA ACGACATTAC TGAACATCAT TCTGGGACAG  1020
GAAACAGCAG AAGGAAGTGT ATGGGTGTCG CCGTCCGCAA ACATCGGCTA TTTAACGCAG  1080
GAGGTGTTTG ATTTGCCTTT AGAACAAACA CCGGAAGAGT TATTTGAGAA TGAAACATTC  1140
AAAGCAAGGG GGCACGTTCA AAATCTGATG AGGCACTTAG GTTTTACAGC CGCCCAATGG  1200
ACTGAACCGA TCAAGCATAT GAGTATGGGT GAGCGTGTAA AGATCAAGCT GATGGCATAT  1260
ATTCTGGAGG AAAAAGACGT GCTGATTTTA GATGAGCCGA CAAACCATCT CGACCTGCCG  1320
TCACGCGAAC AGCTGGAAGA AACACTGTCA CAATACAGCG GCACATTGCT GGCGGTTTCA  1380
CATGACCGAT ACTTTCTCGA AAAAACAACA AACAGTAAAC TCGTCATCTC AAACAACGGC  1440
ATCGAAAAGC AGTTAAACGA CGTTCCTTCA GAAAGAAATG AGCGGGAGGA GCTTCGGTTA  1500
AAGCTTGAGA CAGAAAGACA AGAAGTGCTG GGAAAGCTCA GTTTTATGAC GCCAAATGAT  1560
AAAGGGTATA AGGAGCTTGA TCAGGCTTTC AATGAGCTTA CGAAACGAAT AAAAGAGCTG  1620
GATCATCAAG ACAAAAAAGA CTGA                                         1644
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Figure 4A

MetLysGluIleValThrLeuThrAsnValSerTyrGluValLysAspGlnThrValPhe 20
LysHisValAsnAlaSerValGlnGlnGlyAspIleIleGlyIleIleGlyLysAsnGly 40
AlaGlyLysSerThrLeuLeuHisLeuIleHisAsnAspLeuAlaProAlaGlnGlyGln 60
IleLeuArgLysAspIleLysLeuAlaLeuValGluGlnGluThrAlaAlaTyrSerPhe 80
AlaAspGlnThrProAlaGluLysLysLeuLeuGluLysTrpHisValProLeuArgAsp 100
PheHisGlnLeuSerGlyGlyGluLysLeuLysAlaArgLeuAlaLysGlyLeuSerGlu 120
AspAlaAspLeuLeuLeuLeuAspGluProThrAsnHisLeuAspGluLysSerLeuGln 140
PheLeuIleGlnGlnLeuLysHisTyrAsnGlyThrValIleLeuValSerHisAspArg 160
TyrPheLeuAspGluAlaAlaThrLysIleTrpSerLeuGluAspGlnThrLeuIleGlu 180
PheLysGlyAsnTyrSerGlyTyrMetLysPheArgGluLysLysArgLeuThrGlnGln 200
ArgGluTyrGluLysGlnGlnLysMetValGluArgIleGluAlaGlnMetAsnGlyLeu 220
AlaSerTrpSerGluLysAlaHisAlaGlnSerThrLysLysGluGlyPheLysGluTyr 240
HisArgValLysAlaLysArgThrAspAlaGlnIleLysSerLysGlnLysArgLeuGlu 260
LysGluLeuGluLysAlaLysAlaGluProValThrProGluTyrThrValArgPheSer 280
IleAspThrThrHisLysThrGlyLysArgPheLeuGluValGlnAsnValThrLysAla 300
PheGlyGluArgThrLeuPheLysAsnAlaAsnPheThrIleGlnHisGlyGluLysVal 320

Figure 4B

AlaIleIleGlyProAsnGlySerGlyLysThrThrLeuLeuAsnIleIleLeuGlyGln 340

GluThrAlaGluGlySerValTrpValSerProSerAlaAsnIleGlyTyrLeuThrGln 360

GluValPheAspLeuProLeuGluGlnThrProGluGluLeuPheGluAsnGluThrPhe 380

LysAlaArgGlyHisValGlnAsnLeuMetArgHisLeuGlyPheThrAlaAlaGlnTrp 400

ThrGluProIleLysHisMetSerMetGlyGluArgValLysIleLysLeuMetAlaTyr 420

IleLeuGluGluLysAspValLeuIleLeuAspGluProThrAsnHisLeuAspLeuPro 440

SerArgGluGlnLeuGluGluThrLeuSerGlnTyrSerGlyThrLeuLeuAlaValSer 460

HisAspArgTyrPheLeuGluLysThrThrAsnSerLysLeuValIleSerAsnAsnGly 480

IleGluLysGlnLeuAsnAspValProSerGluArgAsnGluArgGluGluLeuArgLeu 500

LysLeuGluThrGluArgGlnGluValLeuGlyLysLeuSerPheMetThrProAsnAsp 520

LysGlyTyrLysGluLeuAspGlnAlaPheAsnGluLeuThrLysArgIleLysGluLeu 540

AspHisGlnAspLysLysAsp 547